

Appl. No. : Unknown
Filed : Herewith

- the nucleic acid fragment, the sequence of which extends from position -228 to position -451 relative to the ATG codon of the *TaTrxh2* gene;
 - the nucleic acid fragment, the sequence of which extends from position -451 to position -591 relative to the ATG codon of the *TaTrxh2* gene;
 - the nucleic acid fragment, the sequence of which extends from position -83 to position -228 relative to the ATG codon of the *TaTrxh2* gene;
 - the nucleic acid fragment, the sequence of which is that of the first intron of the *TaTrxh2* gene.
3. An expression cassette, comprising a promoter according to Claim 1.
 4. A recombinant vector, comprising at least one promoter according to Claim 1.
 5. A plant cell transformed with at least one promoter according to Claim 1.
 6. A transgenic plant transformed with at least one promoter according to Claim 1.
 7. The transgenic plant according to Claim 6, wherein said plant is a monocotyledon.

Please add the following claim:

9. A method for controlling the expression of a gene of interest in a plant cell comprising using the promoter of Claim 1.

REMARKS

The specification and Claims 1-7 were amended, Claim 8 cancelled, and Claim 9 added in order to place the present application in condition for conventional practice before the United States Patent and Trademark Office. No new matter has been added. Changes to the specification and claims can be seen on a separate page entitled **VERSION WITH MARKINGS TO SHOW CHANGES MADE**. Insertions are underlined and deletions are in **[bold and brackets]**.

Conclusion

Should any issues arise which may delay examination of the present application the Examiner is respectfully invited to contact the under-signed attorney at the telephone number below.

Appl. No. : Unknown
Filed : Herewith

Please charge any additional fees, including any fees for additional extension of time, or credit overpayment to Deposit Account No. 11-1410.

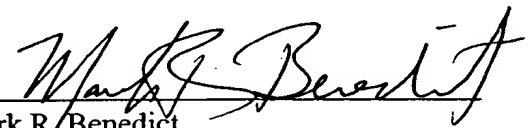
Respectfully submitted,

KNOBBE, MARTENS, OLSON & BEAR, LLP

Dated: _____

11/19/01

By: _____


Mark R. Benedict
Registration No. 44,531
Attorney of Record
620 Newport Center Drive
Sixteenth Floor
Newport Beach, CA 92660
(714) 949-6323

O:\DOCS\CMS\CMS-1984.DOC
111601

Appl. No. : Unknown
Filed : Herewith

VERSION WITH MARKINGS TO SHOW CHANGES MADE

Insertions are underlined and deletions are in **[bold and brackets]**.

IN THE SPECIFICATION:

On page 1, line 2, following the TITLE OF THE INVENTION, please insert the following paragraph:

CROSS REFERENCE TO RELATED APPLICATIONS

This application is a National Phase application under 35 U.S.C. § 371 of PCT application PCT/FR00/01318, filed May 17, 2000 and published in French, which claims priority to French application FR99/06231, filed May 17, 1999. The French application is incorporated herein by reference.

On page 1, line 2, before the paragraph beginning "The invention relates...", please insert the following heading:

FIELD OF THE INVENTION

On page 1, line 3, before the paragraph beginning "Thioredoxins are low...", please insert the following heading:

BACKGROUND OF THE INVENTION

On page 1, line 12, before the paragraph beginning "In plants...", please insert the following heading:

DETAILED DESCRIPTION OF THE INVENTION

On page 23, line 1, please omit the heading "Claims" and insert the following heading:
WHAT IS CLAIMED IS:

IN THE ABSTRACT:

Please insert the abstract attached hereto, following the page entitled VERSION WITH MARKINGS TO SHOW CHANGES MADE, as page 25 of the specification.

IN THE CLAIMS:

Please cancel Claim 8.

Please amend the claims as follows:

1. A p[P]romoter, comprising[characterized in that it consists of] a nucleic acid fragment comprising at least one specific functional domain of the promoter of the *TaTrxh2* gene.
2. The p[P]romoter according to Claim 1, wherein[characterized in that] said nucleic acid fragment is chosen from the group consisting of:
 - the nucleic acid fragment, the sequence of which extends from position -1 to position -1111 relative to the ATG codon of the *TaTrxh2* gene;
 - the nucleic acid fragment, the sequence of which extends from position -1 to position -83 relative to the ATG codon of the *TaTrxh2* gene;
 - the nucleic acid fragment, the sequence of which extends from position -451 to position -591 relative to the ATG codon of the *TaTrxh2* gene;
 - the nucleic acid fragment, the sequence of which extends from position -591 to position -1111 relative to the ATG codon of the *TaTrxh2* gene;
 - the nucleic acid fragment, the sequence of which extends from position -228 to position -451 relative to the ATG codon of the *TaTrxh2* gene;
 - the nucleic acid fragment, the sequence of which extends from position -451 to position -591 relative to the ATG codon of the *TaTrxh2* gene;
 - the nucleic acid fragment, the sequence of which extends from position -83 to position -228 relative to the ATG codon of the *TaTrxh2* gene;
 - the nucleic acid fragment, the sequence of which is that of the first intron of the *TaTrxh2* gene.
3. An e[E]xpression cassette, comprising[characterized in that it comprises] a promoter according to [either of]Claim[s] 1[and 2].
4. A r[R]ecombinant vector, comprising[characterized in that it comprises] at least one promoter according to [either of]Claim[s] 1[and 2].
5. A p[P]lant cell transformed with at least one promoter according to [either of]Claim[s] 1[and 2].

Appl. No. : Unknown
Filed : Herewith

6. A t[T]ransgenic plant transformed with at least one promoter according to [either of
]Claim[s] 1[and 2].

7. The t[T]ransgenic plant according to Claim 6, wherein said plant[characterized in
that it] is a monocotyledon.